

Sparse tensor dimensionality reduction with application to clustering of functional connectivity

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1. ABSTRACT

Functional connectivity (FC) is a graph-like data structure commonly used by neuroscientists to study the dynamic behaviour of the brain activity. However, these analyses rapidly become complex and time-consuming. In this work, we present complementary empirical results on two tensor decomposition previously proposed named modified High Order Orthogonal Iteration (mHOOI) and High Order sparse Singular Value Decomposition (HOsSVD). These decomposition associated to k -means were shown to be useful for the study of multi trial functional connectivity dataset.

Keywords: dynamic networks, graph decomposition, clustering, dimensionality reduction, sparsity, tensor decompositions, HOOI, functional connectivity, iEEG.

2. INTRODUCTION

Epilepsy is one of the most common neurological disorders in the world population. iEEG electrodes¹ are used to exhibit the stages of a seizure distinguished by similar pattern in different areas of the brain. Functional Connectivities (FC) that quantify along time these similarities are calculated between all pairs of signals, usually by means of the spectral coherence or the Phase Locking Value.² Considering electrodes as nodes and FCs as weights on the edges, neuroscientists try to find in the data, graph-like structures evolving through time (see Fig. 1). It is generally assumed that the FC dynamics are comparable from one seizure to another for a same patient, another asset of iEEG monitoring is the possibility to record several seizures. In,³ we proposed a procedure able to extract relevant ensembles of simultaneously activated components, viewed as the edges of sub-graphs of a FC network inferred from the iEEG signals.

More concretely, for each seizure, the recorded iEEG signals yield L pairwise Functional Connectivity measures (FC) as time series of T samples. The corresponding data matrix $\mathbf{X} \in \mathbb{R}^{L \times T}$ is referred to as an Epoch. As we observe S different seizures for a same patient, the resulting epochs are stacked in a 3-modes tensor $\mathcal{X} \in \mathbb{R}^{L \times T \times S}$. The data analysis process, presented Fig. 2, corresponds to first apply a dimensionality reduction method in order to denoise and reduce the dimensionality of the dataset, then apply k -means to the reduction. Notice that directly applying an extension of k -means to the tensor \mathcal{X} corresponds to maximize the following criteria :

$$\operatorname{argmax}_{\mathbf{A}} \quad \|\mathbf{A}^t \mathbf{X}_{(L)}\|_F^2 \quad (1)$$

Where \mathbf{A} corresponds to a normalized cluster assignment matrix⁴ and $\mathbf{X}_{(L)} \in \mathbb{R}^{L \times TS}$ is the mode- L matricization of \mathcal{X} . Such generalisation does not take advantage of the tensor structure of the dataset, moreover k -means is known to perform poorly on high dimension (here TS).⁵ In⁶ we compared and proposed several reduction

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method, transforming the tensor \mathcal{X} to a factor matrix $\mathbf{F} \in \mathbb{R}^{L \times K}$ with $K \ll TS$. The final data analysis process can then be resumed by maximizing the following criteria where \mathbf{A} has the same structure as in (1).

$$\operatorname{argmax}_{\mathbf{A}} \quad \|\mathbf{A}^t \mathbf{F}\|_F^2 \quad (2)$$

The objective of this presentation is to present complementary empirical results on two tensor decomposition methods proposed in⁷ named modified High Order Orthogonal Iteration (mHOOI) and High Order sparse Singular Value Decomposition (HOSVD). Indeed both methods have an original algorithmic implementation without theoretical proof of convergence and convergence speed yet. Also HOSVD which is a sparse tensor decomposition has not been compared to tensor decomposition with similar constraints. After recalling the notation, we present mHOOI, HOSVD and similar decompositions for comparison. Finally the convergence property of HOSVD and mHOOI are exhibited via a model of functional connectivity as well as the performance of HOSVD to decompose a dataset in sparse components via the same model and real epileptic datasets.

3. NOTATIONS

Tensors are denoted with bold case calligraphic letters \mathcal{X} , matrices and vectors are denoted respectively in bold upper-case and lower-case \mathbf{X} , \mathbf{x} , and scalars by lower-case letters x . Notice that l, t, s, k, n will be used as indices, and L, T, S, K, N will be reserved to denote their index upper bounds. Here they correspond respectively to the FC mode, time mode, seizure (or trial) mode, number of factors and number of clusters. Then, $\mathbf{x}_{:t}$, resp. $\mathbf{x}_{l:}$, corresponds to the column t , resp. to the row l , of the matrix $\mathbf{X} \in \mathbb{R}^{L \times T}$. The matrices $\mathbf{X}_{l::}$, $\mathbf{X}_{:t:}$ or $\mathbf{X}_{::s}$ correspond to the slices of the tensor $\mathcal{X} \in \mathbb{R}^{L \times T \times S}$. The slice for each trial $\mathbf{X}_{::s}$ is referred to as an epoch.

The matricization corresponds to the matrix representation of a tensor. The mode- L matricization of \mathcal{X} is noted $\mathbf{X}_{(L)} \in \mathbb{R}^{L \times TS}$. Writing $[\mathbf{A}, \mathbf{B}]$ the concatenation of two matrices \mathbf{A} and \mathbf{B} with the same number of rows, the mode-1 matricization reads:

$$\mathbf{X}_{(L)} = [\mathbf{X}_{::1}, \mathbf{X}_{::2}, \dots, \mathbf{X}_{::S}]. \quad (3)$$

Considering $\mathbf{x}, \mathbf{y} \in \mathbb{R}^L$, $\langle \mathbf{x}, \mathbf{y} \rangle = \sum_{l=1}^L x_l y_l$ is the scalar product between two vectors and $\|\mathbf{x}\|_F = \sqrt{\langle \mathbf{x}, \mathbf{x} \rangle}$ the Frobenius norm. These notations can be extended to D -mode arrays,^{3,4} e.g. $\langle \mathbf{X}, \mathbf{Y} \rangle = \langle \operatorname{vec}(\mathbf{X}), \operatorname{vec}(\mathbf{Y}) \rangle$ or $\|\mathcal{X}\|_F = \|\operatorname{vec}(\mathbf{X}_{(L)})\|_F$. The L_1 -norm for a matrix corresponds to:

$$\|\mathbf{X}\|_1 = \sum_{l=1}^L \sum_{t=1}^T |x_{lt}|. \quad (4)$$

Finally we define the Kronecker product between two matrices, noted $\mathbf{C} = \mathbf{A} \otimes \mathbf{B}$ with $\mathbf{A} \in \mathbb{R}^{L \times T}$, $\mathbf{B} \in \mathbb{R}^{K \times N}$ and $\mathbf{C} \in \mathbb{R}^{LK \times TN}$, as:

$$\mathbf{C} = \begin{pmatrix} a_{11}\mathbf{B} & \dots & a_{1T}\mathbf{B} \\ \vdots & & \vdots \\ a_{L1}\mathbf{B} & \dots & a_{LT}\mathbf{B} \end{pmatrix}. \quad (5)$$

4. REDUCTION METHODS

In order to reduce the tensor data $\mathcal{X} \in \mathbb{R}^{L \times T \times S}$ to a relevant factor matrix $\mathbf{F} \in \mathbb{R}^{L \times K}$, we propose the following decomposition :

$$\mathbf{X}_{(L)} \approx \mathbf{F}(\mathbf{w} \otimes \mathbf{V})^t \quad (6)$$

where $\mathbf{w} \in \mathbb{R}^S$ is a vector performing a weighted average of the S epochs in order to extract common patterns, \mathbf{F} is the factor matrix and $\mathbf{V} \in \mathbb{R}^{T \times K}$ corresponds to the temporal activation profile of the different factors. \mathbf{F} is scaled such that $\|\mathbf{w}\|_F^2 = \|\mathbf{v}_{:k}\|_F^2 = 1$ (to remove scaling indeterminacy and transfer all the energy in our factor

matrix). Without constraints on \mathbf{F} , \mathbf{V} and \mathbf{w} , the High Order Orthogonal Iteration (HOOI) or the proposed approach mHOOI can be used to perform this decomposition. However it as be shown in⁷ that constraining \mathbf{F} to be sparse appears to be important because, in comparison to the case without this constraint, \mathbf{F} becomes closer to a cluster assignment matrix. Moreover, it limits the complexity of clusters by reducing the number of FC they contain. This is important in the context of epileptic data where a large number of FC measurements can be passively implied in a neurological process (during the discharge of the seizure for example). As well it seems relevant to also impose \mathbf{V} to be sparse so as to select specific temporal steps of FC activation, eliminating thus period where there is no common activation of FC clusters. To perform this decomposition we proposed a new algorithm HOsSVD, In a similar way, this be done thanks to a state of the art method like the sparse Tucker decomposition proposed in.⁵

4.1 mHOOI

The proposed modified HOOI (noted mHOOI) is as follows, iterating 2 steps in alternance:

- (A) At iteration $(i + 1)$ we assume $\mathbf{W}_i^{(1)}$ to be known. From this vector we compute the common Epoch corresponding to $\mathbf{X}_{(L)}(\mathbf{W}_i^{(1)} \otimes \mathbf{I}) \in \mathbb{R}^{L \times T}$, with $\mathbf{I} \in \mathbb{R}^{T \times T}$ the identity matrix (it corresponds to the mode- L matricization of the contraction product between the tensor \mathcal{X} and the vector $\mathbf{W}_i^{(1)}$ ³). We then compute the low rank approximations $\mathbf{U}_{i+1}^{(K)}$ and $\mathbf{V}_{i+1}^{(K)}$ as the K first components of the singular value decomposition of $\mathbf{X}_{(L)}(\mathbf{W}_i^{(1)} \otimes \mathbf{I}) = \mathbf{U}_{i+1} \mathbf{\Lambda}_{i+1} \mathbf{V}_{i+1}^t$.
- (B) In order to update $\mathbf{W}_{i+1}^{(1)}$ we first filter the tensor \mathcal{X} by projecting it on the subspaces spanned by the matrices $\mathbf{U}_{i+1}^{(K)}$ and $\mathbf{V}_{i+1}^{(K)}$. $\mathbf{W}_{i+1}^{(1)}$ is then obtained by a special HOOI decomposition on the filtered tensor with $K_L = L$, $K_T = T$ and $K_S = 1$. The optimal decomposition can be found analytically. Both the filtering and the decomposition, in order to get $\mathbf{W}_{i+1}^{(1)}$, reduce to the problem of computing the dominant left singular vector of $\mathbf{X}_{(S)}(\mathbf{V}_{i+1}^{(K)} \otimes \mathbf{U}_{i+1}^{(K)})$.

We summarize in Algorithm 1, this modified HOOI method (mHOOI) and how to obtain the corresponding lower dimension factor matrix \mathbf{F} . We use an angular metric between two successive estimates of $\mathbf{W}^{(1)}$ as the stopping criterion of the iterative procedure.

4.2 HOsSVD

The previous algorithm requires several SVD and the final matrix F is not sparse in general. As we already noticed, sparsity constraints is wanted to get FC graphs with few edges, and because the SVD is impacted by the curse of dimensionality it is desirable to limit its use. Following the same rationale, and in order to perform a decomposition as proposed in Eq. (6), we develop a sparse version of mHOOI, where we replace the computation of the SVD in step (A), by the sparse SVD :

$$\underset{\mathbf{F}, \mathbf{Z}}{\operatorname{argmin}} \quad \|\mathbf{X}_{(L)} - \mathbf{F}\mathbf{Z}^t\|_F^2 + \gamma_1 \|\mathbf{F}\|_1, \quad (7)$$

$$s.t. \quad \gamma_2 \|\mathbf{Z}_{:k}\|_1 + \|\mathbf{Z}_{:k}\|_F^2 \leq 1 \quad (8)$$

where the meta-parameters γ_1 and γ_2 allow for tuning the trade-off between accuracy and sparsity of the approximation. To solve this matrix factorization (??), we use the SPAMS library on matlab.⁶ We call this decomposition a High Order sparse SVD (HOsSVD). Since the matrix \mathbf{U}_{i+1} is not necessarily orthogonal anymore, we compute the subspace spanned by this matrix by performing its QR decomposition, and we retain only the first K column vectors of the \mathbf{Q} part. The same goes for \mathbf{V}_{i+1} . Writing $qr(\mathbf{V}_{i+1}, K)$ this operation, Algorithm 2 presents the computation of \mathbf{F} , the lower dimension factor matrix stemming from this high order sparse SVD reduction (HOsSVD).

4.3 State of the art methods

Both proposed approach mHOOI and HOSVD will be compared by state of the art approach. Notice the following decomposition :

$$\underset{\mathbf{U}, \mathbf{V}, \mathbf{W}}{\operatorname{argmin}} \|\mathbf{X}_{(L)} - \mathbf{U}\mathbf{G}_{(L)}(\mathbf{W} \otimes \mathbf{V})^t\|_F^2, \quad (9)$$

Where $\mathbf{U} \in \mathbb{R}^{L \times K_L}$, $\mathbf{V} \in \mathbb{R}^{T \times K_T}$, $\mathbf{W} \in \mathbb{R}^{L \times K_S}$ and $\mathbf{G}_{(L)} \in \mathbb{R}^{K_L \times K_T \times K_S}$ is a dense matrix. The High order orthogonal Iteration (HOOI) algorithm provide a solution to this optimisation problem (eq. 9).^{7,8} Now by noting $\mathbf{F} = \mathbf{U}\mathbf{G}_{(L)}$ in the case $K_L = K$, $K_T = K$ and $K_S = 1$ we get a solution of the decomposition eq. 6 similar to the mHOOI algorithm 1.

In,⁵ they propose a decomposition which constraints matrices \mathbf{U} , \mathbf{V} , \mathbf{W} and $\mathbf{G}_{(L)}$ to be non negative and optionally sparse. By fixing $K_L = K$, $K_T = K$ and $K_S = 1$ and imposing sparsity on \mathbf{U} and \mathbf{V} we get a decomposition similar to HOSVD. The factor matrix will be $\mathbf{F} = \mathbf{U}$ after scaling \mathbf{U} such as $\|\mathbf{w}\|_F^2 = \|\mathbf{v}_{:k}\|_F^2 = \mathbf{G}_{(L)} : k = 1$ for all $k \in 1, \dots, K$. This decomposition will is called SN-Tucker.

Algorithm 1 Estimation of \mathbf{F} via mHOOI

Require: \mathcal{X} , K , angular tolerance $\epsilon > 0$, et i_{max} .

```

i = 0
 $\mathbf{W}_0^{(1)} = \text{left-1-SVD}(\mathbf{X}_{(3)})$ 
while i < imax or  $\operatorname{acos}(\langle \mathbf{W}_i^{(1)}, \mathbf{W}_{i-1}^{(1)} \rangle) > \epsilon$  do
  (A).  $[\mathbf{U}_{i+1}^{(K)}, \mathbf{\Lambda}_{i+1}, \mathbf{V}_{i+1}^{(K)}] = K\text{-SVD}(\mathbf{X}_{(L)}(\mathbf{W}_i^{(1)} \otimes \mathbf{I}))$ 
  (B).  $\mathbf{W}_{i+1}^{(1)} = \text{left-1-SVD}(\mathbf{X}_{(S)}(\mathbf{V}_{i+1}^{(K)} \otimes \mathbf{U}_{i+1}^{(K)}))$ 
  i = i + 1
end while
 $\mathbf{F} = \mathbf{U}_i^{(K)} \mathbf{\Lambda}_i$ 

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Algorithm 2 Estimation of \mathbf{F} via HOSVD

Require: \mathcal{X} , K , γ_1 , γ_2 , angular tolerance $\epsilon > 0$, et

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imax.
i = 0
 $\mathbf{W}_0^{(1)} = \text{left-1-SVD}(\mathbf{X}_{(3)})$ 
while i < imax or  $\operatorname{acos}(\langle \mathbf{W}_i^{(1)}, \mathbf{W}_{i-1}^{(1)} \rangle) > \epsilon$  do
  (A).  $\mathbf{U}_{i+1}, \mathbf{V}_{i+1}$  minimising (??) with
   $\mathbf{X}_{(L)}(\mathbf{W}_i^{(1)} \otimes \mathbf{I})$ .
  (B).  $\mathbf{W}_{i+1}^{(1)} = \text{left-1-SVD}(\mathbf{X}_{(S)}(\mathbf{B} \otimes \mathbf{A}))$ 
  i = i + 1
end while
 $\mathbf{F} = \mathbf{U}_i^{(K)}$ 

```

5. RESULTS AND APPLICATIONS

In this section we present different empirical results on the convergence speed of mHOOI, the convergence of HOSVD and his performances on a model of FC. First, we present the model, then the different empirical results. Finally the HOSVD method is applied on real epileptic data.

Matlab code for HOOI, mHOOI, and HOSVD can be found here ^{*}. These codes uses the MATLAB Tensor toolbox Version 2.6 [†] and the sparse SVD is done using the SPAMS toolbox version 2.6 [‡]? Finally the SN-Tucker algorithm come from the toolbox [§]

5.1 Model

Epileptic seizure implies a pathological FC that starts in a focal brain onset, then spreads to the other connected regions, and sometimes split to give rise to new FC components. Fig. 3 displays a characteristic example of actual FC time series measured by the phase lock value (PLV).² The model we propose is aimed at reproducing the global structured pattern of FCs' activation, and the uncertainties of the measures. More precisely, we consider a matrix $\mathbf{X} \in \mathbb{R}^{L \times T}$, where x_{lt} is set to a high value if the FC of index $l \in \{1, \dots, L\}$ is active at time $t \in \{1, \dots, T\}$, and to a low value otherwise (to account for the non ON-OFF discrepancy of the PLV measurement, we choose values equal to 0.7 and 0.2, respectively). A cluster C_n , $n \in \{1, \dots, N\}$, is composed of all FCs that are activated over the same period of time T_n , figure 4 present clusters of FC activation. This is for the deterministic part of

^{*}Algorithms used in this article: FCTensDec[?]

[†]MATLAB Tensor Toolbox Version 2.6³

[‡]SPAMS toolbox version 2.6⁶

[§]www2.imm.dtu.dk/pubdb/views/edoc_download.php/4718/zip/imm4718.zip⁵

the model, defining the structural and temporal pattern of FCs activation, common to all seizures. Superimposed to it, we add a seizure dependant random component, composed of four uncertainty sources:

- i Random duration: each activation period T_n is uniformly distributed between a minimum duration (here, 5 time steps) and $\frac{T}{N}$ time steps (the binary variable α allows to able (1) or disable (0) this random mode).
- ii Activation error: with probability $\beta \in [0, 1]$, each FC of a given group C_n incurs the risk to be replaced by any other randomly chosen FC.
- iii Connectivity noise: we add to x_{lt} a Uniform noise in the interval $[0, \sigma]$ if the FC is inactivated, $[1 - \sigma, 1]$ else.
- iv Jitter: All FCs of the same group start activating with independent jitters, uniformly distributed in $[-\frac{\delta}{2}, \frac{\delta}{2}]$.

Figure 5 displays one realization of these synthetic FC time series. Compared to real data, the model succeeds in reproducing a realistic global pattern. More importantly, as it allows a control of the nature and the intensity of variability between epochs, it will serve to evaluate the sensitivity and the robustness of the different dimension reduction methods for clustering, with respect to each source of uncertainty.

To this end, we simulate different seizures of a same patient as i.i.d. realizations of our model with the same set of parameters $b = [\alpha, \beta, \sigma, \delta]$. As for real data, the S modeled seizures are then stacked in a tensor $\mathcal{X} \in \mathbb{R}^{L \times T \times S}$. Figure 6 displays 4 realisations of the model $\mathbf{X}_{:,i}$ corresponding to the particular choice $[\alpha = 1, \beta = 0.2, \sigma = 0.6, \delta = 0.3]$, $L = 66$, $T = 100$ and $S = 4$.

5.2 On the convergence speed of mHOOI vs HOOI

To empirically observe the convergence towards the optimal solution, with faster performance than HOOI algorithm we propose the following experiment. We consider tensors from the model presented above, with different configuration of noise vector. We fix $K_L = K_T = 3$ and $K_S = 1$. We compare the performance of both HOOI and mHOOI algorithm by computing the total variance at each iteration $tvar(t) = \|\mathbf{G}_i\|_F^2$ for HOOI, $tvar(t) = \|\mathbf{F}_i\|_F^2$ (since they are matrices containing all the variance) and the log differential of the total variance between two consecutive iterates $ldvar(t) = \log(tvar(t) - tvar(t - 1))$. Fig. 7 shows the mean of $tvar(t)$ for the first 100 iterations using 100 realisations of the tensor model \mathcal{X} associated tot he noise vector b_i as input of the algorithm. A zoom of the last iteration is provided as well as the first value of $ldvar(t)$ is shown. Notice the lake of value of $ldvar$ is due to the fact that in average $tvar(t) - tvar(t - 1)$ is negative, however it is not relevant since it only happen when this difference attains the precision of the machine. The new proposed algorithm has always better performances than HOOI, with exponential convergence, on every scenario. (??change fig add b_i ??)

5.3 On the convergence of HOsSVD

As previously we consider the model presented above, we study the convergence of the HOsSVD alone. Since the objective of SN-Tucker and HOsSVD are not exactly the same it is useless to compare them like mHOOI and HOOI.

For a set of hyper parameters γ_1 and γ_2 we compute the first 100 iterations using 100 realisations of the tensor model \mathcal{X} associated to the noise vector b_i as input of the HOsSVD algorithm (2). We say that the algorithm converge on 100 iteration if the angular tolerance criterion is attain at the 100th iteration or before. The table 1 shows for different noise vector b_i and hyper parameters the percentage over 120 experiments that converge on 100 iterations. Few situations lead to potential no convergence of HOsSVD algorithm: cases b_2 with $\gamma_1 = 1$, $\gamma_2 = 1$ and b_9 with $\gamma_1 = 1$ converge $\approx 90\%$ of the time. There is two kind of non convergence, the first kind happening once for b_9 with $\gamma_1 = 1$, $\gamma_2 = 1$ is due to high sparsity constraints that lead to decomposition in null matrices. The second kind of non convergence happen more frequently, where the algorithm oscillates between several solutions. The figure 8 shows the convergence criteria for 12 realisations with the noise vector b_{10} and $\gamma_1 = 1$, $\gamma_2 = 1$, where we can observe the second kind of non convergence. Notice convergence always happen where there is connectivity noise (which is always the case, with different probability distribution, in real dataset).

Finally in order to show that the convergence of this algorithm is fast, the table 1 shows for different noise vector b_i and hyper parameters the percentage over 120 experiments that converge on 5 iterations. We observe fast

convergence when there is connectivity noise in the model. The figure 9 shows the logarithm of the convergence criteria for the same 12 realisations as the figure 8 highlighting the exponential convergence of the algorithm as the mHOOI algorithm (notice the algorithm is stopped when the angular tolerance criteria is attained, explaining the stationary phase).

5.4 On the Performances of HOsSVD vs SN-Tucker

5.4.1 Experimental setup

Using the FC model, we compare HOsSVD and SN-Tucker methods derived in Section ???. Each method yields a lower dimension factor matrix \mathbf{F} that serves as an input to k -means. Performance refer here to the ability at retrieving the FC clusters of the global pattern (fig. (4)). We use a set of hyper-parameters values to compare both method, notice the effect of these parameters does not impact both methods equally, however they are chosen to cover the principal behaviours of these algorithms. We fix K to 3 since there is 3 activated FC clusters in the model. Regarding k -means algorithm, we used k -means++ version⁹ that we stopped after 1000 iterations and repeated 120 times with different seeds. The number of cluster is set to $N = 4$. Finally, to assess the clustering performance of each method, we use the Adjusted Rand Index (ARI) score,¹⁰ computed between the resulting grouping and the ground truth figure 4: the score equals 1 for a perfect match and 0 if the correspondence does not outperform a random grouping.

5.4.2 Result

Under these experimental conditions, we evaluate all methods for various configurations of uncertainty and hyper-parameters. Fig. 10 displays the ARI scores (mean and variance estimated out of 120 independent realizations of \mathcal{X}) for 10 different combinations of (i) random duration, (ii) activation error and (iii) connectivity and (iv) jitter noise. For each noise configuration, we highlight the hyper-parameters which give the best performance for both method by a horizontal line.

HOsSVD outperform SN-Tucker for all configurations considering the best performances of each methods along the set of hyper-parameters. Also we can observe more variability of the ARI score using HOsSVD methods with different hyper parameters, showing a stronger impact of the hyper parameters on the final result. in fact, for high constraints of sparsity, our proposed approach seems to give more relevant decompositions than SN-Tucker, indeed it is less likely to get identical factors using HOsSVD than SN-Tucker. This can be shown figure 11 where for a realisation on the noise configuration $b = [0, 0, 0.7, 0.3]$ we show the factor matrix F for every hyper parameters using HOsSVD method, we can see for high constraints of sparsity non equivalent factors and a result close to an assignment cluster matrix. Contrary to the previous case, the factor matrix F obtained using SN-Tucker (fig. (??)) method seems not relevant, with null factors for high constraints of sparsity.

5.5 Application on real data

5.5.1 Data

We consider real iEEG data from 3 patients with focal epilepsy.^{1,11} The electrodes used are distributed on stems implanted in the brain. The activity of the brain is recorded via 5 to 10 electrodes per stem. The space between two consecutive electrodes is 3.5 mm. Over a recording time of 15 days, 4 seizures were selected for the first patient, 4 for the second patient and 2 for the third. We only have clinical knowledge of the seizure for the first patient. Each seizure is delimited in time by a window of 100 seconds centered on the beginning of the seizures. The signal is sampled at 256 Hz. Only a thirds of the initial contacts are selected to avoid too strong spatial correlations. The functional connectivity metric used is PLV.² A strong PLV between two signals means that their phases are similar. The FC (corresponding to each pair of electrodes) were calculated over a sliding rectangular window of 4 seconds duration, with a time step of one second. After eliminating the points that suffer from border effects, the data is formatted as a tensor $\mathcal{X} \in \mathbb{R}^{528 \times 96 \times 4}$ for the first patient, $\mathcal{X} \in \mathbb{R}^{1035 \times 96 \times 4}$ for the second and $\mathcal{X} \in \mathbb{R}^{946 \times 96 \times 2}$ for the third. The FC of each seizure are plotted in 13, 14 and 15 respectively for the patient 1, 2 and 3.

5.5.2 Methods

From the tensor \mathcal{X} , we obtain the matrix \mathbf{F} (using Algorithm 2), where the parameters, $K = 4$, $\epsilon = 10^{-3}$ and $\gamma_1 = 1$, $\gamma_2 = 1$ for the first patient, $\gamma_1 = 0.8$, $\gamma_2 = 1.6$ for the second patient, $\gamma_1 = 1.2$, $\gamma_2 = 0.8$ for the third patient are empirically fixed (It is current to consider between 3 and 5 different steps in an epileptic seizure justifying $K = 4$; for γ_1 and γ_2 we search for the highest value of parameters where results seem coherent). Fig. 16 17 18 (top) shows the temporal activation profiles of the components of \mathbf{F} (corresponding to the matrix \mathbf{V} from Algorithm 2). We apply k -means to \mathbf{F} to identify the $N = 5$ corresponding FC groups. Since \mathbf{F} is close to an assignment matrix (a solution \mathbf{A}^* of Eq. (2)), we retain the 4 groups of smaller sizes, which can be associated with the 4 activation periods while the 5th group corresponds to the unsynchronized FCs. Figure 16 17 18 (bottom) materializes the positions of the 33 electrodes projected on the transverse plane (according to the lair of Tailarach) for each patient. An FC is represented by a link between the pair of electrodes that are in phase.

5.5.3 Results

Patient 1 with 4 epileptic seizure and clinical knowledge 16: There are 4 activation periods that can be easily associated with 4 steps of the seizures: before seizure, start, propagation and end of the seizure. The time interval around time 50s is particularly interesting as it shows no activated FC. This is likely to correspond to the functional decoupling at the early start of the seizure, a short period when iEEG activities in different areas of the brain are suddenly decorrelated. The four FC groups can be associated to four snapshots of a time-varying graph: before the seizure, only two electrodes interact in what could correspond to the epileptogenic zone. At the beginning of the seizure: spreading of FC activation with the appearance of a cluster of FC localized around the epileptogenic zone. During the crisis, other FC appears spontaneously in the other hemisphere, at the same time the FC are diffused in the left hemisphere. At the end of the crisis, the two hemispheres interact with the appearance of common FC. The graphs obtained are in agreement with the clinical results, the focus of the beginning of the seizure being close to the graphs "before-seizure" and "seizure start". The propagation of seizures in the right hemisphere of the brain is well represented by the graphs "Propagation" and "Seizure end".

Patient 2 with 4 epileptic seizure without clinical knowledge 17: As well as the previous patient 4 activation periods that can be easily associated with 4 steps of the seizures: before seizure, start, propagation and end of the seizure. There is more period without activation, certainly because FC patterns of each seizure seems more erratic than the previous patient. The four FC groups can aswell be associated to four snapshots of a time-varying graph: at step 1 only two electrodes interact, at step 2 and 3 spreading of FC activation in all the brain. The last step present we return to a situation close to the first.

Patient 3 with 2 epileptic seizure without clinical knowledge 18: Contrary to the two previous seizure, it is more difficult to distinct activation period, this can be explained by the fact that we only have access to two epileptic seizure for this patient, which mean less robust results. However the graph corresponding to each activation are localized spatially, which indicate a cohenrency in the obtained results.

6. CONCLUSION

We have presented new tensor decompositions with original algorithmic implementation. We shown through a general model of FC the fast empirical convergence of these algorithms compared to the already existing HOOI. The advantage of using HOsVD instead of a state of the art tensor non negative sparse tensor decomposition to find sparse FC common to each seizure is exhibited. Finally, applied to real iEEG data recorded during an epileptic seizure, our method allowed us to identify FC activation groups corresponding to significant time periods of the evolution of the seizure.

As a perspective of this work, we would like to generalise the proposed tensor reduction to the general case where the seizure mode can be decomposed in several factor ($K_S > 1$).

7. ACKNOWLEDGEMENT

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Matlab code from⁷ including HOSVD and mHOOI algorithm can be found here ⁴:

⁴FCTensDec

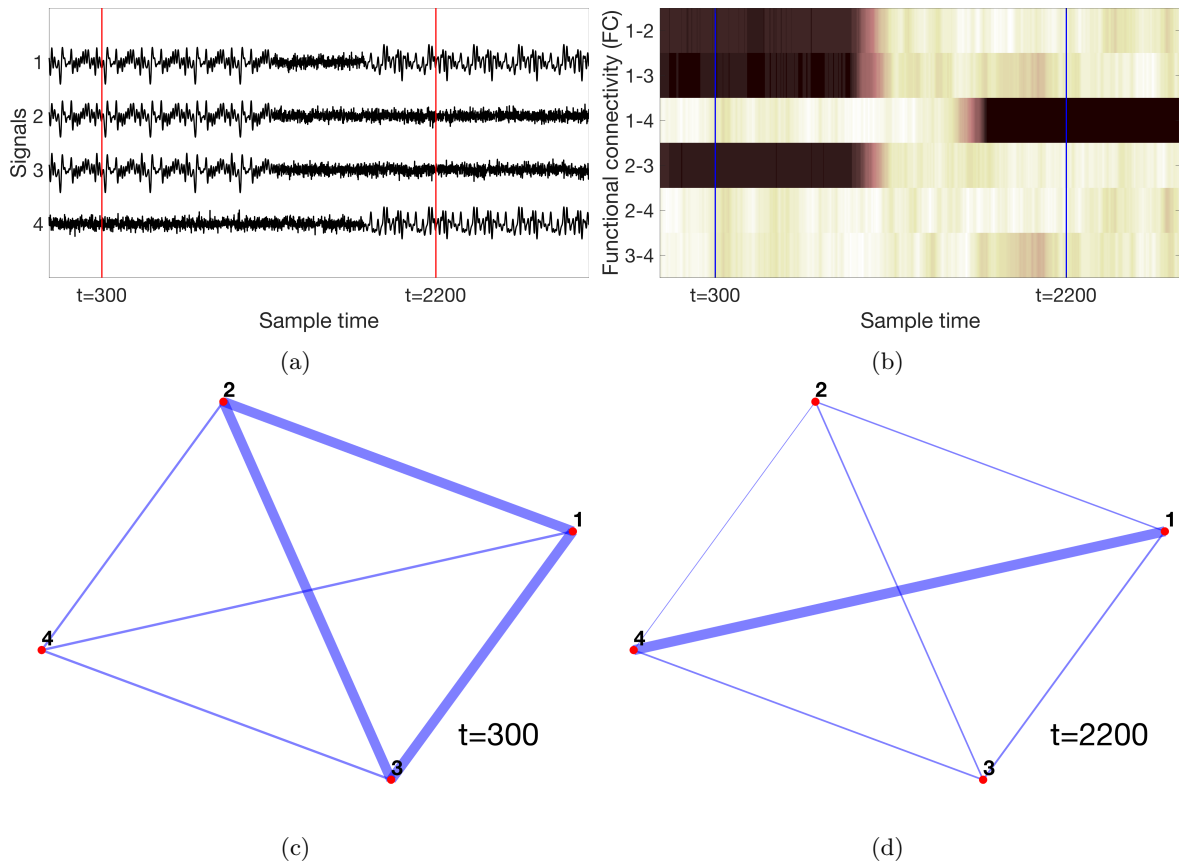


Figure 1: (a) 4 simulated iEEG signals, (b) FC computed between the 6 pairs of signals by PLV, (c) & (d) Graphs with electrodes 1,2,3 and 4 as nodes and PLV measures as the weight of edges, at $t=300$ and $t=2200$ (red/blue bars on Fig. (a) and (b)).

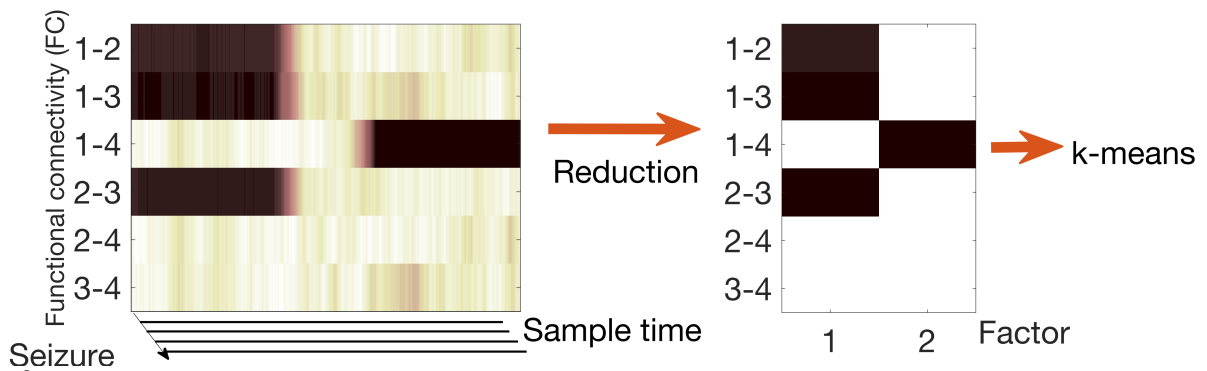


Figure 2: Model of the data processing pipeline: tensor \mathcal{X} is reduced to a factor matrix \mathbf{F} , and k -means is applied on \mathbf{F}

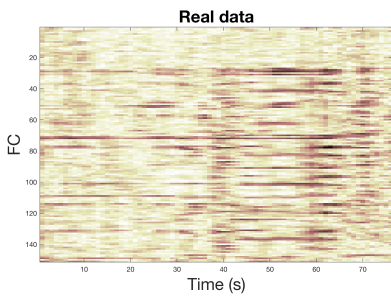


Figure 3

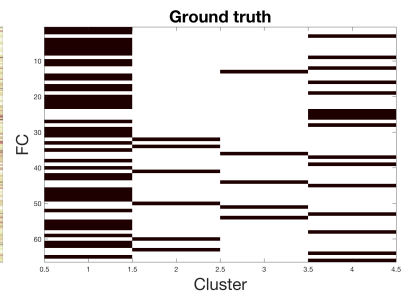


Figure 4

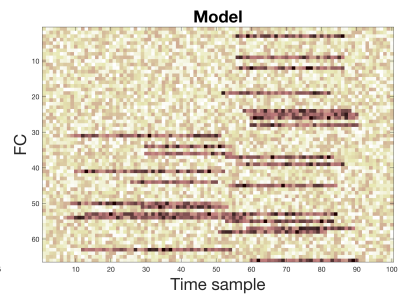
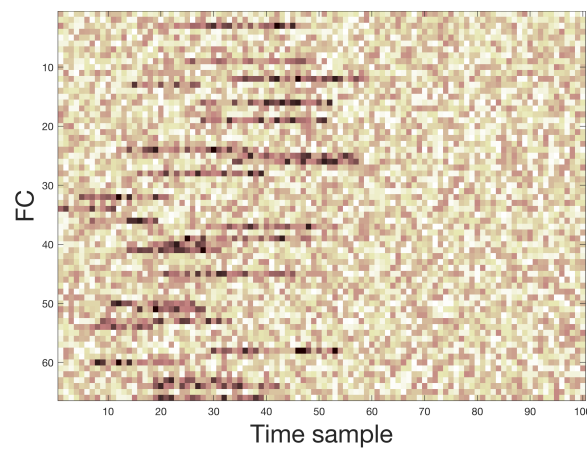
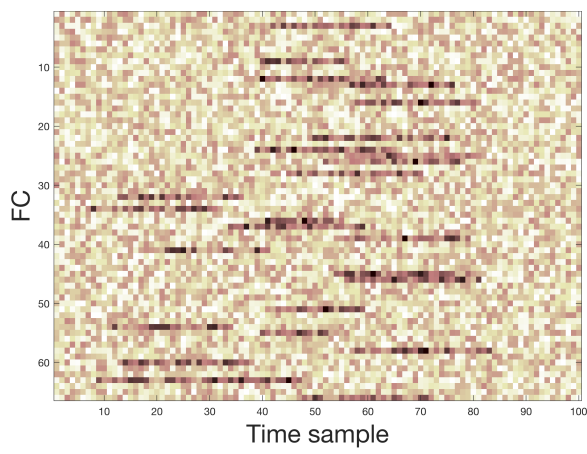


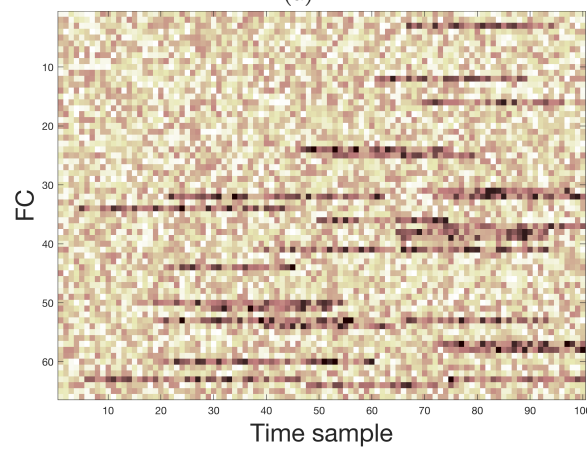
Figure 5



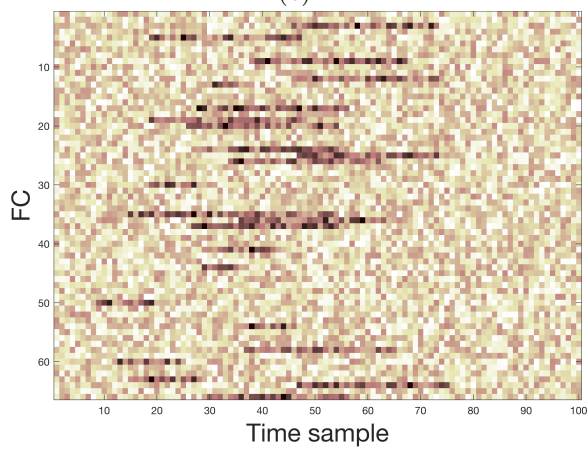
(a)



(b)



(c)



(d)

Figure 6: Example of 4 realisation our model with $b = [1, 0.2, 0.6, 0.3]$

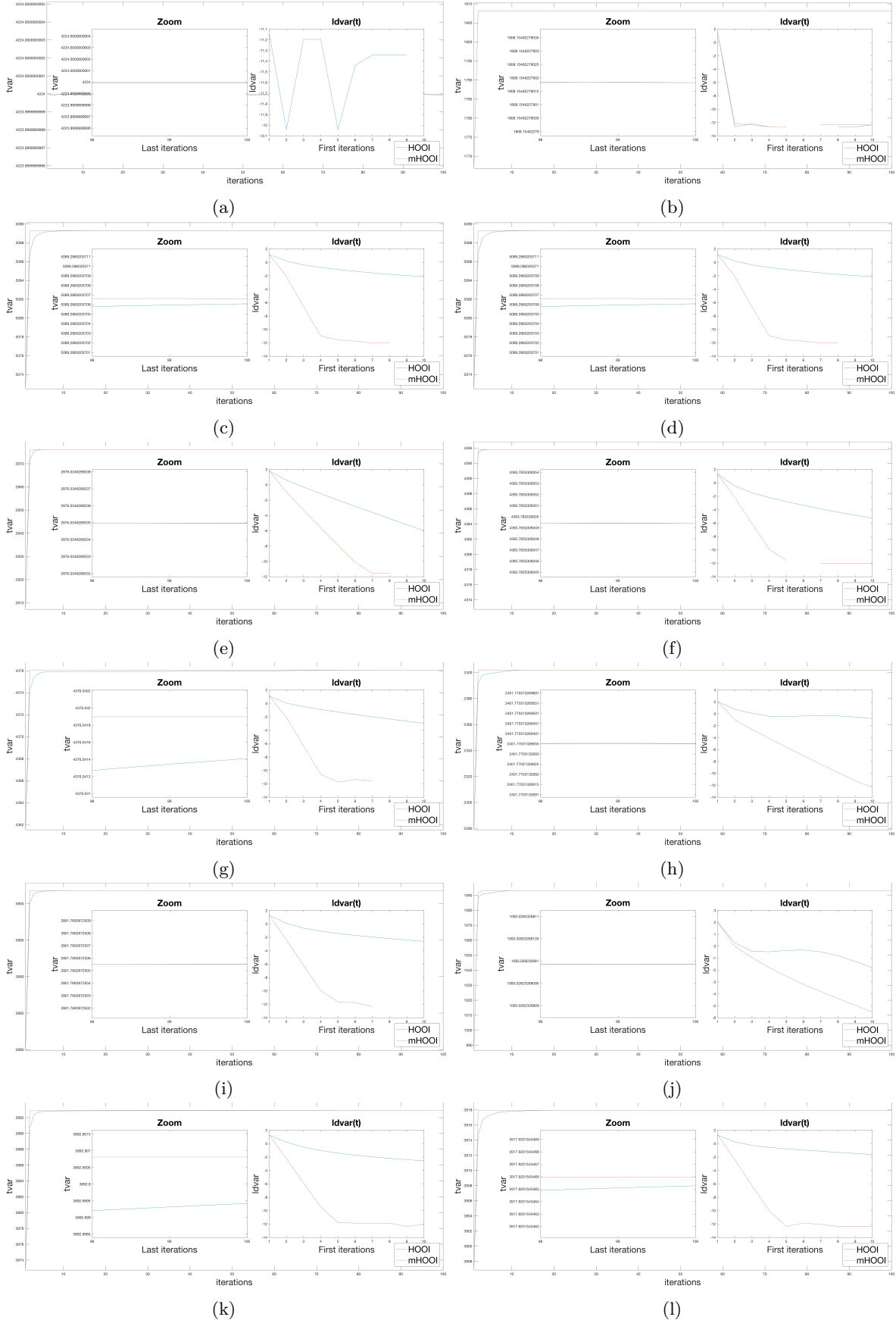


Figure 7: $tvar(t)$ for the first 100 iterations using the tensor \mathcal{X}_{model} as input of the algorithm with a zoom of the last iteration and the first value of $ldvar(t)$ where (a) corresponds to the configuration b_1 , (b) b_2 , (c) b_3 , (d) b_4 , (e) b_5 , (f) b_6 , (g) b_8 , (h) b_8 , (i) b_9 , (j) b_{10} , (k) b_{11} , (l) b_{12} ,

$[\gamma_1, \gamma_2] =$	[0.1, 0.1]	[0.1, 2]	[2, 0.1]	[0.4, 0.4]	[0.4, 1]	[1, 0.4]	[0.7, 0.7]	[1, 0.7]	[0.7, 1]	[1, 1]
$b_1 = [0, 0, 0, 0]$	100	100	100	100	100	100	100	100	100	100
$b_2 = [1, 0, 0, 0]$	100	100	100	100	100	100	100	100	100	91.6667
$b_3 = [0, 0.3, 0, 0]$	100	100	100	100	100	100	100	100	100	100
$b_4 = [0, 0, 0.8, 0]$	100	100	100	100	100	100	100	100	100	100
$b_5 = [0, 0, 0, 0.4]$	100	100	100	100	100	100	100	100	100	100
$b_6 = [0, 0.2, 0.7, 0]$	100	100	100	100	100	100	100	100	100	100
$b_7 = [0, 0, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	100
$b_8 = [0, 0.2, 0, 0.3]$	100	100	100	100	100	100	100	100	100	100
$b_9 = [1, 0.2, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	100
$b_{10} = [1, 0.2, 0, 0.3]$	100	100	100	100	100	91.6667	100	91.6667	100	91.6667
$b_{11} = [1, 0.2, 0.7, 0]$	100	100	100	100	100	100	100	100	100	100
$b_{12} = [1, 0, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	100

Table 1: % of 120 experiment which converge at least before **100** iteration using HOSVD algorithm with parameters γ_1, γ_2 and with the noise configuration b_i .

	[0.1, 0.1]	[0.1, 2]	[2, 0.1]	[0.4, 0.4]	[0.4, 1]	[1, 0.4]	[0.7, 0.7]	[1, 0.7]	[0.7, 1]	[1, 1]
$b_1 = [0, 0, 0, 0]$	100	100	100	100	100	100	100	100	100	100
$b_2 = [1, 0, 0, 0]$	83.3333	33.3333	33.3333	33.3333	8.3333	8.3333	8.3333	8.3333	8.3333	0
$b_3 = [0, 0.3, 0, 0]$	83.3333	0	0	0	0	8.3333	0	0	0	0
$b_4 = [0, 0, 0.8, 0]$	100	100	100	100	100	100	100	100	100	100
$b_5 = [0, 0, 0, 0.4]$	100	100	100	100	100	100	100	91.6667	91.6667	75
$b_6 = [0, 0.2, 0.7, 0]$	100	100	100	100	100	100	100	91.6667	100	83.3333
$b_7 = [0, 0, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	100
$b_8 = [0, 0.2, 0, 0.3]$	50	33.3333	25	33.3333	8.3333	33.3333	33.3333	25	25	8.3333
$b_9 = [1, 0.2, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	8.3333
$b_{10} = [1, 0.2, 0, 0.3]$	0	0	0	0	0	0	0	0	0	0
$b_{11} = [1, 0.2, 0.7, 0]$	100	100	100	100	100	100	100	75	75	16.6667
$b_{12} = [1, 0, 0.7, 0.3]$	100	100	100	100	100	100	100	100	100	25

Table 2: % of 120 experiment which converge at least before **5** iteration using HOSVD algorithm with parameters γ_1, γ_2 and with the noise configuration b_i .

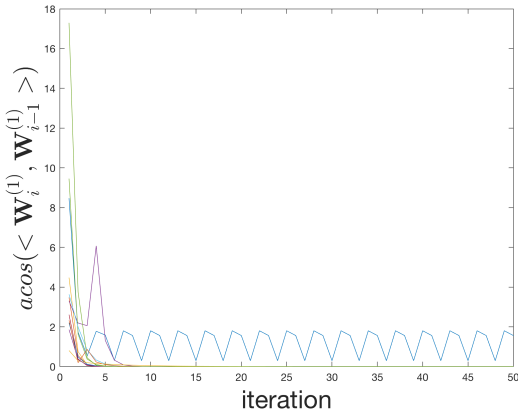


Figure 8: Convergence criteria for 12 realisation the realisation of the experiment configuration b_{10} and experiment configuration b_{10} and $[\gamma_1, \gamma_2] = [1, 1]$. $[\gamma_1, \gamma_2] = [1, 1]$.

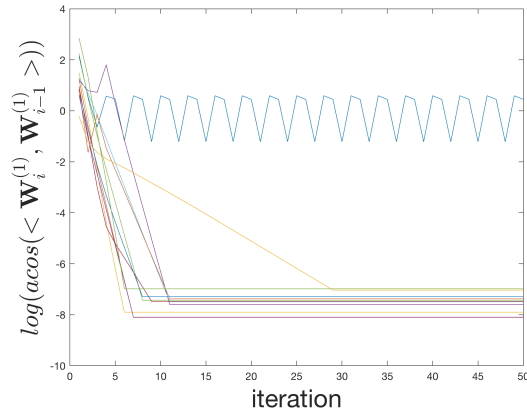


Figure 9: Log of the convergence criteria for 12

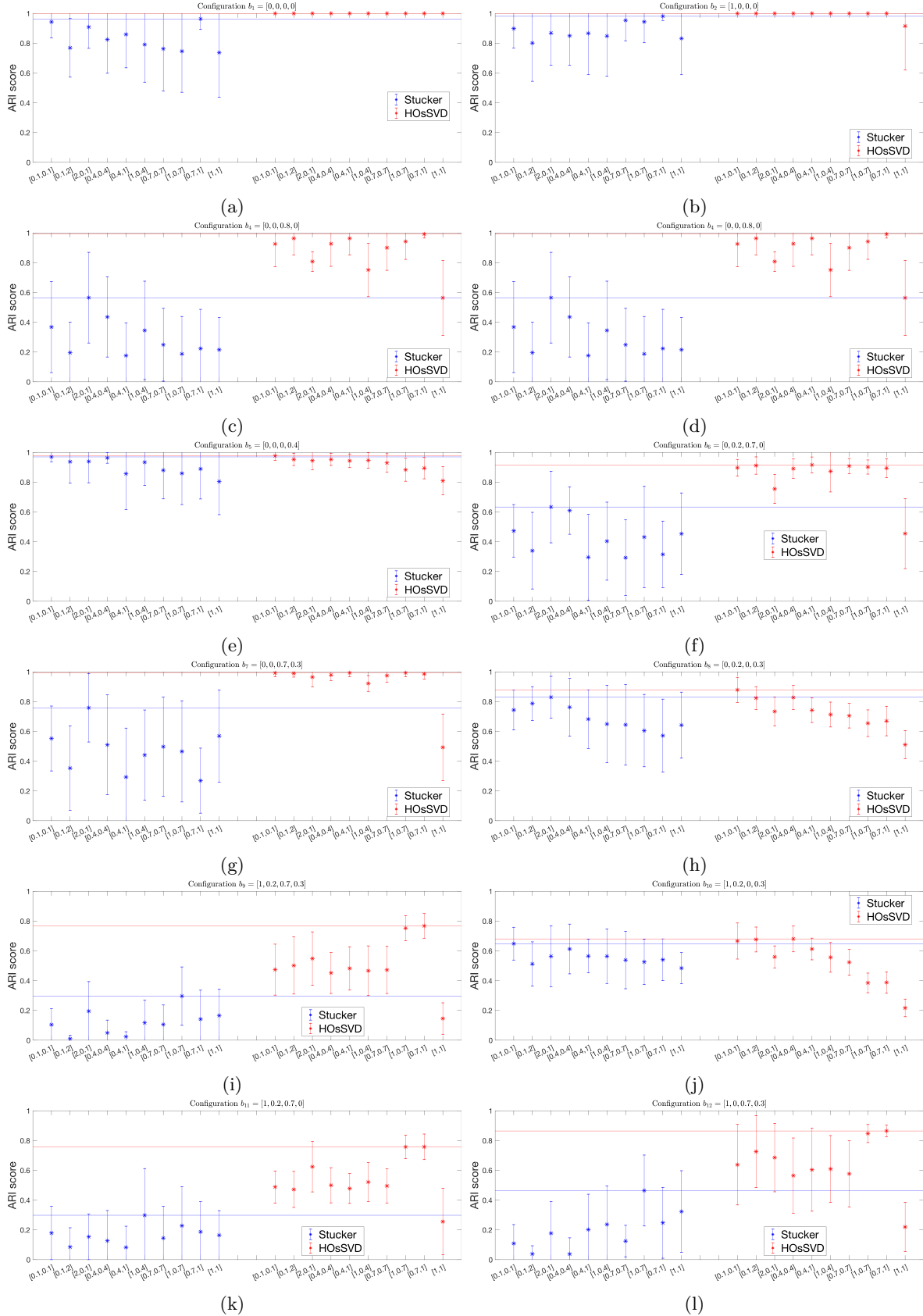


Figure 10: ARI score using SN-Tucker and HO-SVD algorithm in function of the sparsity parameters and the configurations

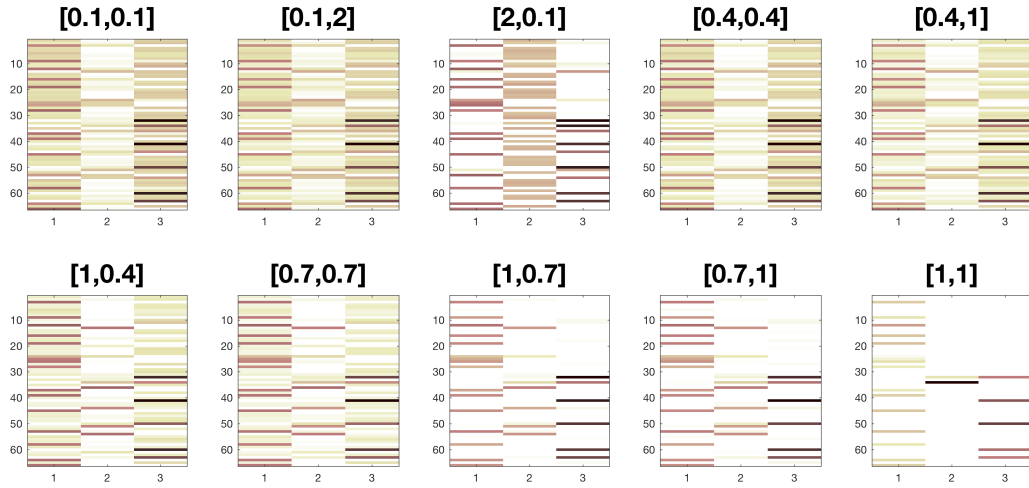


Figure 11: F reduction for all sparsity configurations using HOSVD

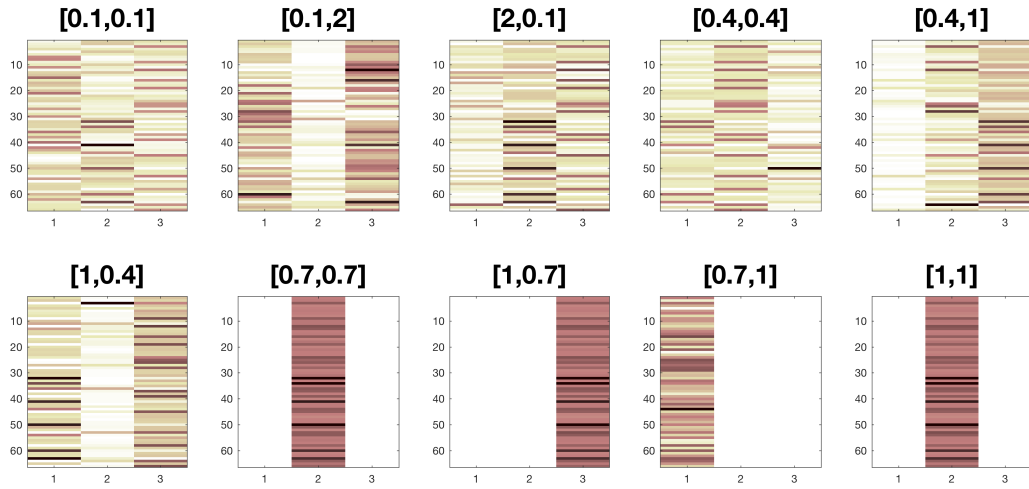
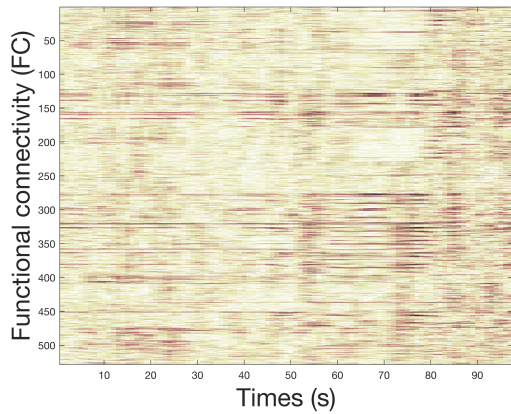
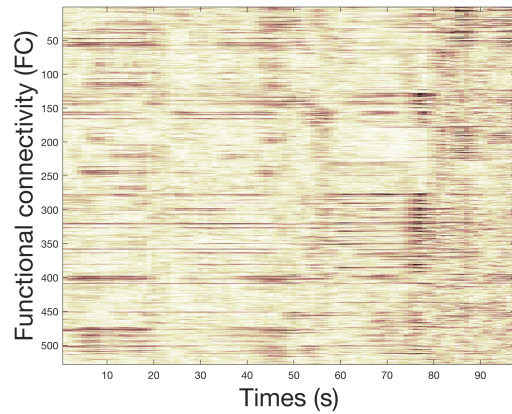


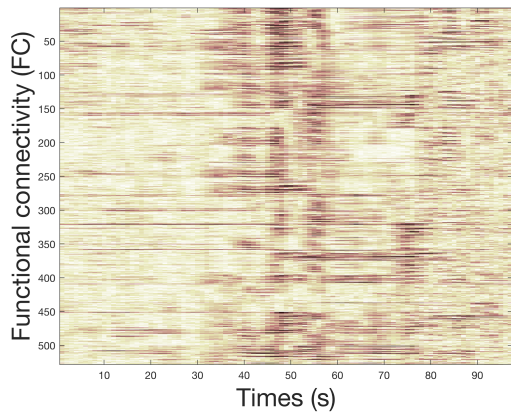
Figure 12: F reduction for all sparsity configurations using SN-Tucker



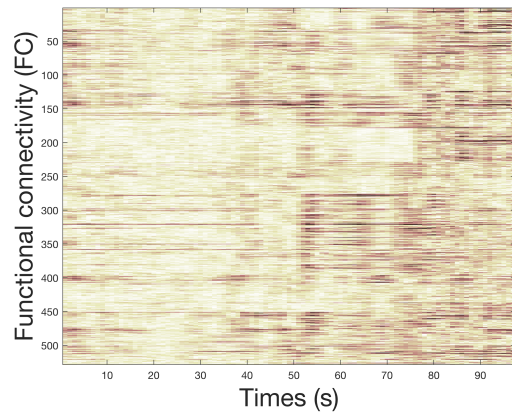
(a)



(b)

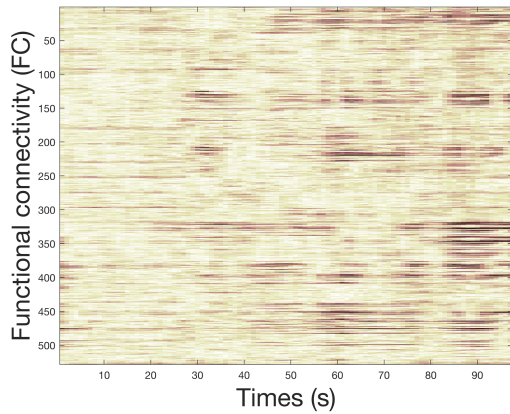


(c)

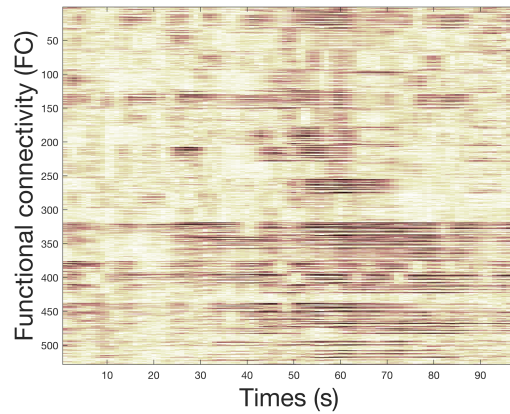


(d)

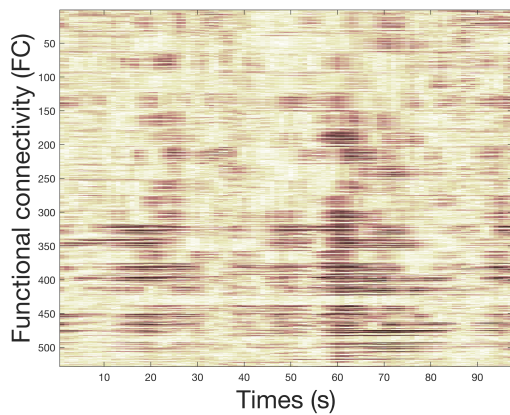
Figure 13: Functional connectivity of all 4 seizures for the patient 1



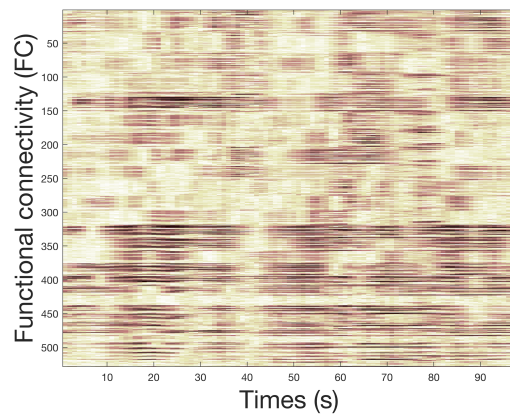
(a)



(b)

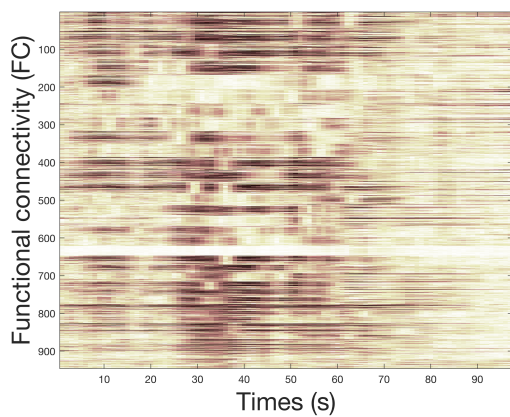


(c)

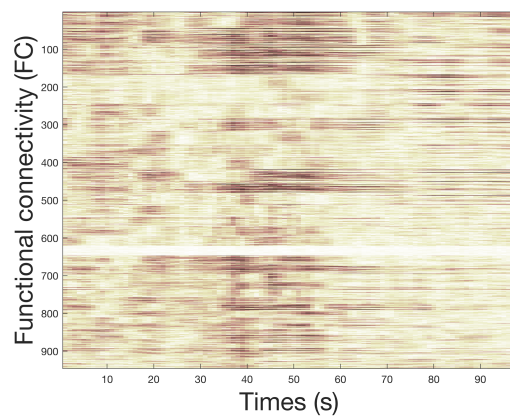


(d)

Figure 14: Functional connectivity of all 4 seizures for the patient 2



(a)



(b)

Figure 15: Functional connectivity of the 2 seizures for the patient 3

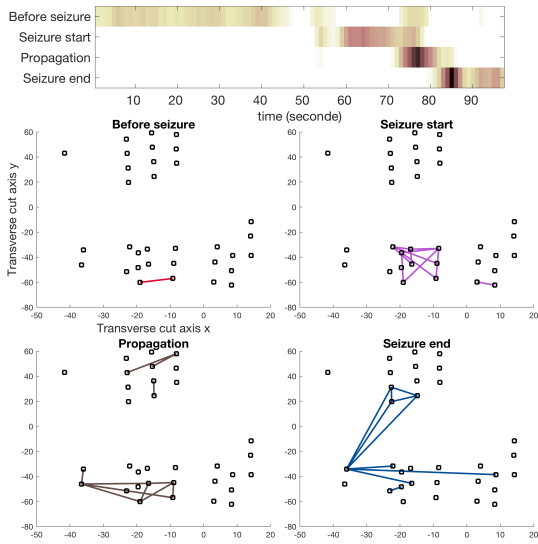


Figure 16: patient 1) : (above) activation profile of HOsVD (variable \mathbf{V} of algorithm 2 (under) Cluster of FC corresponding to the 4 activation steps of the seizures

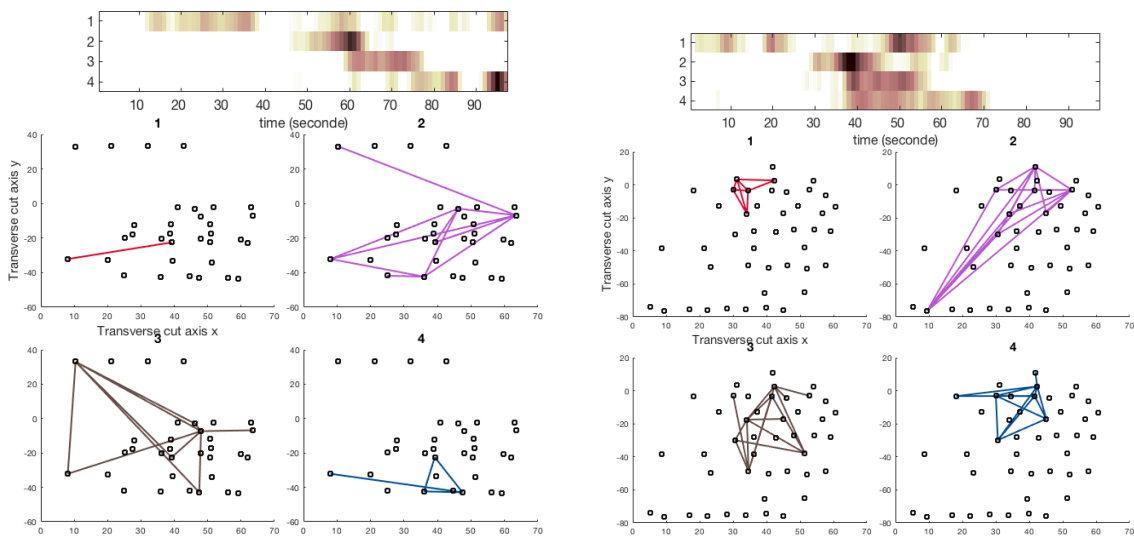


Figure 17: patient 2) : (above) activation profile of HOsVD (variable \mathbf{V} of algorithm 2 (under) Cluster of FC corresponding to the 4 activation steps of the seizures.

REFERENCES

- [1] Guenot, M., Isnard, J., Ryvlin, P., Fischer, C., Ostrowsky, K., Mauguiere, F., and Sindou, M., “Neurophysiological monitoring for epilepsy surgery: the talairach seeg method,” *Stereotactic and functional neurosurgery* **77**(1-4), 29–32 (2001).
- [2] van Mierlo, P., Papadopoulou, M., Carrette, E., Boon, P., Vandenberghe, S., Vonck, K., and Marinazzo, D., “Functional brain connectivity from EEG in epilepsy: Seizure prediction and epileptogenic focus localization,” *Progress in neurobiology* **121**, 19–35 (2014).

- [3] Kolda, T. G. and Bader, B. W., “Tensor decompositions and applications,” *SIAM review* **51**(3), 455–500 (2009).
- [4] De Lathauwer, L., De Moor, B., and Vandewalle, J., “A multilinear singular value decomposition,” *SIAM journal on Matrix Analysis and Applications* **21**(4), 1253–1278 (2000).
- [5] Mørup, M., Hansen, L. K., and Arnfred, S. M., “Algorithms for sparse nonnegative tucker decompositions,” *Neural computation* **20**(8), 2112–2131 (2008).
- [6] Mairal, J., Bach, F., Ponce, J., Sapiro, G., Jenatton, R., and Obozinski, G., “SPAMS: A SPArse Modeling Software, v2. 3,” *URL <http://spams-devel.gforge.inria.fr/downloads.html>* (2014).
- [7] Zhang, A. and Xia, D., “Tensor SVD: Statistical and Computational Limits,” *IEEE Transactions on Information Theory* (2018).
- [8] De Lathauwer, L., De Moor, B., and Vandewalle, J., “On the best rank-1 and rank-(r_1, r_2, \dots, r_n) approximation of higher-order tensors,” *SIAM journal on Matrix Analysis and Applications* **21**(4), 1324–1342 (2000).
- [9] Arthur, D. and Vassilvitskii, S., “k-means++: The advantages of careful seeding,” in [*Proceedings of the eighteenth annual ACM-SIAM symposium on Discrete algorithms*], 1027–1035, Society for Industrial and Applied Mathematics (2007).
- [10] Hubert, L. and Arabie, P., “Comparing partitions,” *Journal of classification* **2**(1), 193–218 (1985).
- [11] Chauvel, P., Rheims, S., McGonigal, A., and Kahane, P., “French guidelines on stereoelectroencephalography (seeg): Editorial comment.,” *Neurophysiologie clinique= Clinical neurophysiology* **48**(1), 1 (2018).